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methylation ("whole genome amplification") (c) Advanced Scholar Search

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In vitro DNA immortalization and whole genome amplification using libraries generated from randomly fragmented DNA

PH Pister, T Kurnikova, I Slepakova ... - US Patent App 10/1 ... 2004 - Google Patents
 ... (43) Pub. Date: Oct 21, 2004 (54) IN VITRO DNA IMMORTALIZATION AND WHOLE GENOME AMPLIFICATION USING LIBRARIES GENERATED FROM RANDOMLY FRAGMENTED DNA Related US Application Data (60) Provisional application No. 60/453,071, filed on Mar. ...

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The Hematopoietic System-specific Minor Histocompatibility Antigen HA-1 Shows Aberrant Expression in Epithelial Cancer Cells

CA Hwang, M J Kim, J H Kim, ... - The Journal of ... 2002 - jpmexpress.org
 ... HA-1 CD45 and PBGD expression were quantitated in all test samples using these calibration functions. The isolated DNA was subjected to whole genome amplification and CSH. ... DNA methylation has been proposed as the primary mechanism for the inactivation of genes (24). ...

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[HTML] from jpmexpress.org

Multiplexed Methylation Detection Methods

JB Jain - US Patent App 10/731,203 ... 2003 - Google Patents
 ... Date: Nov. 22, 2007 (54) MULTIPLEXED METHYLATION DETECTION METHODS (76) Inventor: Jian-Bing Fan, San Diego, CA (US) Correspondence Address: MCDERMOTT, WILL & EMERY 4370 LA JOLLA VILLAGE DRIVE, SUITE 700 SAN DIEGO, CA 92122 (US) (21) Appl. ...

Multiplexed Methylation Detection Methods

JB Jain - US Patent App 20,070,284,601, 2003 - freepatentsonline.com
 ... The invention also provides a calibration curve comprising a first axis indicating the amount of unmethylated template population compared to methylated template ... The invention methods for detecting methylation allow for whole genome amplification after bisulfite ...

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The San Diego Conference Nucleic Acid Technology: The Cutting Edge of Discovery November 6-8, 1997

S. Graham - Critical Care Medicine 1997 - Am Assoc Crit Care ...
 ... To address these limitations, we have combined whole genome amplification by primer extension ... the non-competitive nature of the amplification of the target and quantitation standard. ... Regulatory region mutations as well as methylation represent additional possible mechanisms ...

Polynucleotide sequence detection assays

BB Rosenblum - US Patent App. 201,650,93,857, 2006 - Google Patents
 ... addition, preparation of target nucleic acids can be accomplished using whole genome amplification techniques (eg ... also be used to estimate the average amount of methylation for one ... by including nucleotide analogs having higher binding affinities than standard nucleotides for ...

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Polynucleotide sequence detection assays

BB Rosenblum - US Patent App. 107,18,016, 2003 - Google Patents
 ... of target nucleic acids can be accomplished using whole genome amplification techniques (eg. Lizardi, US Pat. No. 6,124,120). [0041] In some embodiments, target nucleic acids are chemically treated prior to analysis. For example, analysis of the methylation state of cytosines ...

Behavioral and metabolic consequences of neonatal exposure to diazepam in rat pups

H Schroeder, AC Humber, V Kozel, D Desir ... - Experimental ... 1985 - Elsevier
 ... for 15 ... *Developmental ...* 20, Diazep ... Ad. J. Dev. Res.

Cellular engineering, protein expression profiling, differential labeling of peptides, and novel reagents therefor

JM Smart, M Lattimore, J Wei ... - US Patent App. 20,1,620 ... 2002 - freepatentsonline.com
 ... labeling reagents which relied on stable isotopes, which are expensive, and not flexible to differential labeling of more than two mixtures of peptides; labeling methods limited only to methylation of carboxy ... Standard IUPAC nomenclature is used for describing pooled nucleotides ...

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methylation ("whole genome amplif[

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